

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/508,343
Source: P4710
Date Processed by STIC: 3/13/05

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PCT

RAW SEQUENCE LISTING

DATE: 03/13/2005

PATENT APPLICATION: US/10/508,343

TIME: 11:58:40

Input Set : A:\US10508343-seq list.txt

Output Set: N:\CRF4\03132005\J508343.raw

3 <110> APPLICANT: Evotec NeuroSciences GmbH
 5 <120> TITLE OF INVENTION: DIAGNOSTIC AND THERAPEUTIC USE OF HUMAN MAGUIN PROTEINS
 6 AND NUCLEIC ACIDS FOR NEURODEGENERATIVE DISEASES
 8 <130> FILE REFERENCE: 020791ep ME/BM
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/508,343
 C--> 11 <141> CURRENT FILING DATE: 2004-09-20
 13 <160> NUMBER OF SEQ ID NOS: 24
 15 <170> SOFTWARE: PatentIn Ver. 2.1
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 1034
 19 <212> TYPE: PRT
 20 <213> ORGANISM: Homo sapiens
 22 <400> SEQUENCE: 1
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 24 1 5 10 15
 26 Val Asp Trp Met Lys Gly Leu Asp Asp Cys Leu Gln Gln Tyr Ile Lys
 27 20 25 30
 29 Asn Phe Glu Arg Glu Lys Ile Ser Gly Asp Gln Leu Leu Arg Ile Thr
 30 35 40 45
 32 His Gln Glu Leu Glu Asp Leu Gly Val Ser Arg Ile Gly His Gln Glu
 33 50 55 60
 35 Leu Ile Leu Glu Ala Val Asp Leu Leu Cys Ala Leu Asn Tyr Gly Leu
 36 65 70 75 80
 38 Glu Thr Glu Asn Leu Lys Thr Leu Ser His Lys Leu Asn Ala Ser Ala
 39 85 90 95
 41 Lys Asn Leu Gln Asn Phe Ile Thr Gly Arg Arg Arg Ser Gly His Tyr
 42 100 105 110
 44 Asp Gly Arg Thr Ser Arg Lys Leu Pro Asn Asp Phe Leu Thr Ser Val
 45 115 120 125
 47 Val Asp Leu Ile Gly Ala Ala Lys Ser Leu Leu Ala Trp Leu Asp Arg
 48 130 135 140
 50 Ser Pro Phe Ala Ala Val Thr Asp Tyr Ser Val Thr Arg Asn Asn Val
 51 145 150 155 160
 53 Ile Gln Leu Cys Leu Glu Leu Thr Thr Ile Val Gln Gln Asp Cys Thr
 54 165 170 175
 56 Val Tyr Glu Thr Glu Asn Lys Ile Leu His Val Cys Lys Thr Leu Ser
 57 180 185 190
 59 Gly Val Cys Asp His Ile Ile Ser Leu Ser Ser Asp Pro Leu Val Ser
 60 195 200 205
 62 Gln Ser Ala His Leu Glu Val Ile Gln Leu Ala Asn Ile Lys Pro Ser
 63 210 215 220
 65 Glu Gly Leu Gly Met Tyr Ile Lys Ser Thr Tyr Asp Gly Leu His Val
 66 225 230 235 240

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68 Ile Thr Gly Thr Thr Glu Asn Ser Pro Ala Asp Arg Cys Lys Lys Ile
69                245                250                255
71 His Ala Gly Asp Glu Val Ile Gln Val Asn His Gln Thr Val Val Gly
72                260                265                270
74 Trp Gln Leu Lys Asn Leu Val Asn Ala Leu Arg Glu Asp Pro Ser Gly
75                275                280                285
77 Val Ile Leu Thr Leu Lys Lys Arg Pro Gln Ser Met Leu Thr Ser Ala
78                290                295                300
80 Pro Ala Leu Leu Lys Asn Met Arg Trp Lys Pro Leu Ala Leu Gln Pro
81 305                310                315                320
83 Leu Ile Pro Arg Ser Pro Thr Ser Ser Val Ala Thr Pro Ser Ser Thr
84                325                330                335
86 Ile Ser Thr Pro Thr Lys Arg Asp Ser Ser Ala Leu Gln Asp Leu Tyr
87                340                345                350
89 Ile Pro Pro Pro Pro Ala Glu Pro Tyr Ile Pro Arg Asp Glu Lys Gly
90                355                360                365
92 Asn Leu Pro Cys Glu Asp Leu Arg Gly His Met Val Gly Lys Pro Val
93                370                375                380
95 His Lys Gly Ser Glu Ser Pro Asn Ser Phe Leu Asp Gln Glu Tyr Arg
96 385                390                395                400
98 Lys Arg Phe Asn Ile Val Glu Glu Asp Thr Val Leu Tyr Cys Tyr Glu
99                405                410                415
101 Tyr Glu Lys Gly Arg Ser Ser Ser Gln Gly Arg Arg Glu Ser Thr Pro
102                420                425                430
104 Thr Tyr Gly Lys Leu Arg Pro Ile Ser Met Pro Val Glu Tyr Asn Trp
105                435                440                445
107 Val Gly Asp Tyr Glu Asp Pro Asn Lys Met Lys Arg Asp Ser Arg Arg
108                450                455                460
110 Glu Asn Ser Leu Leu Arg Tyr Met Ser Asn Glu Lys Ile Ala Gln Glu
111 465                470                475                480
113 Glu Tyr Met Phe Gln Arg Asn Ser Lys Lys Asp Thr Gly Lys Lys Ser
114                485                490                495
116 Lys Lys Lys Gly Asp Lys Ser Asn Ser Pro Thr His Tyr Ser Leu Leu
117                500                505                510
119 Pro Ser Leu Gln Met Asp Ala Leu Arg Gln Asp Ile Met Gly Thr Pro
120                515                520                525
122 Val Pro Glu Thr Thr Leu Tyr His Thr Phe Gln Gln Ser Ser Leu Gln
123                530                535                540
125 His Lys Ser Lys Lys Lys Asn Lys Gly Pro Ile Ala Gly Lys Ser Lys
126 545                550                555                560
128 Arg Arg Ile Ser Cys Lys Asp Leu Gly Arg Gly Asp Cys Glu Gly Trp
129                565                570                575
131 Leu Trp Lys Lys Lys Asp Ala Lys Ser Tyr Phe Ser Gln Lys Trp Lys
132                580                585                590
134 Lys Tyr Trp Phe Val Leu Lys Asp Ala Ser Leu Tyr Trp Tyr Ile Asn
135                595                600                605
137 Glu Glu Asp Glu Lys Ala Glu Gly Phe Ile Ser Leu Pro Glu Phe Lys
138                610                615                620
140 Ile Asp Arg Ala Ser Glu Cys Arg Lys Lys Tyr Ala Phe Lys Ala Cys

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141 625          630          635          640
143 His Pro Lys Ile Lys Ser Phe Tyr Phe Ala Ala Glu His Leu Asp Asp
144          645          650          655
146 Met Asn Arg Trp Leu Asn Arg Ile Asn Met Leu Thr Ala Gly Tyr Ala
147          660          665          670
149 Glu Arg Glu Arg Ile Lys Gln Glu Gln Asp Tyr Trp Ser Glu Ser Asp
150          675          680          685
152 Lys Glu Glu Ala Asp Thr Pro Ser Thr Pro Lys Gln Asp Ser Pro Pro
153          690          695          700
155 Pro Pro Tyr Asp Thr Tyr Pro Arg Pro Pro Ser Met Ser Cys Ala Ser
156 705          710          715          720
158 Pro Tyr Val Glu Ala Lys His Ser Arg Leu Ser Ser Thr Glu Thr Ser
159          725          730          735
161 Gln Ser Gln Ser Ser His Glu Glu Phe Arg Gln Glu Val Thr Gly Ser
162          740          745          750
164 Ser Ala Val Ser Pro Ile Arg Lys Thr Ala Ser Gln Arg Arg Ser Trp
165          755          760          765
167 Gln Asp Leu Ile Glu Thr Pro Leu Thr Ser Ser Gly Leu His Tyr Leu
168          770          775          780
170 Gln Thr Leu Pro Leu Glu Asp Ser Val Phe Ser Asp Ser Ala Ala Ile
171 785          790          795          800
173 Ser Pro Glu His Arg Arg Gln Ser Thr Leu Pro Thr Gln Lys Cys His
174          805          810          815
176 Leu Gln Asp His Tyr Gly Pro Tyr Pro Leu Ala Glu Ser Glu Arg Met
177          820          825          830
179 Gln Val Leu Asn Gly Asn Gly Gly Lys Pro Arg Ser Phe Thr Leu Pro
180          835          840          845
182 Arg Asp Ser Gly Phe Asn His Cys Cys Leu Asn Ala Pro Val Ser Ala
183          850          855          860
185 Cys Asp Pro Gln Asp Asp Val Gln Pro Pro Glu Val Glu Glu Glu Glu
186 865          870          875          880
188 Glu Glu Glu Glu Glu Glu Gly Glu Ala Ala Gly Glu Asn Ile Gly Glu
189          885          890          895
191 Lys Ser Glu Ser Arg Glu Glu Lys Leu Gly Asp Ser Leu Gln Asp Leu
192          900          905          910
194 Tyr Arg Ala Leu Glu Gln Ala Ser Leu Ser Pro Leu Gly Glu His Arg
195          915          920          925
197 Ile Ser Thr Lys Met Glu Tyr Lys Leu Ser Phe Ile Lys Arg Cys Asn
198          930          935          940
200 Asp Pro Val Met Asn Glu Lys Leu His Arg Leu Arg Ile Leu Lys Ser
201 945          950          955          960
203 Thr Leu Lys Ala Arg Glu Gly Glu Val Ala Ile Ile Asp Lys Val Leu
204          965          970          975
206 Asp Asn Pro Asp Leu Thr Ser Lys Glu Phe Gln Gln Trp Lys Gln Met
207          980          985          990
209 Tyr Leu Asp Leu Phe Leu Asp Ile Cys Gln Asn Thr Thr Ser Asn Asp
210          995          1000          1005
212 Pro Leu Ser Ile Ser Ser Glu Val Asp Val Ile Thr Ser Ser Leu Ala
213          1010          1015          1020

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215	His	Thr	His	Ser	Tyr	Ile	Glu	Thr	His	Val							
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219	<210>	SEQ	ID	NO:	2												
220	<211>	LENGTH:	948														
221	<212>	TYPE:	PRT														
222	<213>	ORGANISM:	Homo sapiens														
224	<400>	SEQUENCE:	2														
225	Phe	Ile	Gly	Arg	Glu	Ser	Glu	Gln	Ile	Asp	Asn	Ala	Met	Ile	Asn	Ala	
226	1				5					10					15		
228	Cys	Ile	Asp	Ser	Glu	Gln	Glu	Asn	Cys	Glu	Phe	His	Met	Ala	Asn	Met	
229	-	-	-	20	-	-	-	-	25					30			
231	Ala	Gly	Ile	Asn	Pro	Arg	Thr	Glu	Ile	Asn	Leu	Glu	Asn	Gly	Thr	His	
232			35					40					45				
234	Ala	Ala	Met	Ala	Leu	Ile	Met	Glu	Pro	Val	Ser	Lys	Trp	Ser	Pro	Ser	
235	50						55					60					
237	Gln	Val	Val	Asp	Trp	Met	Lys	Gly	Leu	Asp	Asp	Cys	Leu	Gln	Gln	Tyr	
238	65					70				75						80	
240	Ile	Lys	Asn	Phe	Glu	Arg	Glu	Lys	Ile	Ser	Gly	Asp	Gln	Leu	Leu	Arg	
241					85					90					95		
243	Ile	Thr	His	Gln	Glu	Leu	Glu	Asp	Leu	Gly	Val	Ser	Arg	Ile	Gly	His	
244				100					105					110			
246	Gln	Glu	Leu	Ile	Leu	Glu	Ala	Val	Asp	Leu	Leu	Cys	Ala	Leu	Asn	Tyr	
247			115					120				125					
249	Gly	Leu	Glu	Thr	Glu	Asn	Leu	Lys	Thr	Leu	Ser	His	Lys	Leu	Asn	Ala	
250	130					135				140							
252	Ser	Ala	Lys	Asn	Leu	Gln	Asn	Phe	Ile	Thr	Gly	Arg	Arg	Arg	Ser	Gly	
253	145					150				155					160		
255	His	Tyr	Asp	Gly	Arg	Thr	Ser	Arg	Lys	Leu	Pro	Asn	Asp	Phe	Leu	Thr	
256					165				170					175			
258	Ser	Val	Val	Asp	Leu	Ile	Gly	Ala	Ala	Lys	Ser	Leu	Leu	Ala	Trp	Leu	
259				180					185					190			
261	Asp	Arg	Ser	Pro	Phe	Ala	Ala	Val	Thr	Asp	Tyr	Ser	Val	Thr	Arg	Asn	
262			195					200					205				
264	Asn	Val	Ile	Gln	Leu	Cys	Leu	Glu	Leu	Thr	Thr	Ile	Val	Gln	Gln	Asp	
265	210						215					220					
267	Cys	Thr	Val	Tyr	Glu	Thr	Glu	Asn	Lys	Ile	Leu	His	Val	Cys	Lys	Thr	
268	225					230				235						240	
270	Leu	Ser	Gly	Val	Cys	Asp	His	Ile	Ile	Ser	Leu	Ser	Ser	Asp	Pro	Leu	
271					245					250					255		
273	Val	Ser	Gln	Ser	Ala	His	Leu	Glu	Val	Ile	Gln	Leu	Ala				

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288 Ser Gly Val Ile Leu Thr Leu Lys Lys Arg Pro Gln Ser Met Leu Thr
289          340          345          350
291 Ser Ala Pro Ala Leu Leu Lys Asn Met Arg Trp Lys Pro Leu Ala Leu
292          355          360          365
294 Gln Pro Leu Ile Pro Arg Ser Pro Thr Ser Ser Val Ala Thr Pro Ser
295          370          375          380
297 Ser Thr Ile Ser Thr Pro Thr Lys Arg Asp Ser Ser Ala Leu Gln Asp
298 385          390          395          400
300 Leu Tyr Ile Pro Pro Pro Pro Ala Glu Pro Tyr Ile Pro Arg Asp Glu
301          405          410          415
303 Lys Gly Asn Leu Pro Cys Glu Asp Leu Arg Gly His Met Val Gly Lys
304          420          425          430
306 Pro Val His Lys Gly Ser Glu Ser Pro Asn Ser Phe Leu Asp Gln Glu
307          435          440          445
309 Tyr Arg Lys Arg Phe Asn Ile Val Glu Glu Asp Thr Val Leu Tyr Cys
310          450          455          460
312 Tyr Glu Tyr Glu Lys Gly Arg Ser Ser Ser Gln Gly Arg Arg Glu Ser
313 465          470          475          480
315 Thr Pro Thr Tyr Gly Lys Leu Arg Pro Ile Ser Met Pro Val Glu Tyr
316          485          490          495
318 Asn Trp Val Gly Asp Tyr Glu Asp Pro Asn Lys Met Lys Arg Asp Ser
319          500          505          510
321 Arg Arg Glu Asn Ser Leu Leu Arg Tyr Met Ser Asn Glu Lys Ile Ala
322          515          520          525
324 Gln Glu Glu Tyr Met Phe Gln Arg Asn Ser Lys Lys Asp Thr Gly Lys
325          530          535          540
327 Lys Ser Lys Lys Lys Gly Asp Lys Ser Asn Ser Pro Thr His Tyr Ser
328 545          550          555          560
330 Leu Leu Pro Ser Leu Gln Met Asp Ala Leu Arg Gln Asp Ile Met Gly
331          565          570          575
333 Thr Pro Val Pro Glu Thr Thr Leu Tyr His Thr Phe Gln Gln Ser Ser
334          580          585          590
336 Leu Gln His Lys Ser Lys Lys Lys Asn Lys Gly Pro Ile Ala Gly Lys
337          595          600          605
339 Ser Lys Arg Arg Ile Ser Cys Lys Asp Leu Gly Arg Gly Asp Cys Glu
340          610          615          620
342 Gly Trp Leu Trp Lys Lys Lys Asp Ala Lys Ser Tyr Phe Ser Gln Lys
343 625          630          635          640
345 Trp Lys Lys Tyr Trp Phe Val Leu Lys Asp Ala Ser Leu Tyr Trp Tyr
346          645          650          655
348 Ile Asn Glu Glu Asp Glu Lys Ala Glu Gly Phe Ile Ser Leu Pro Glu
349          660          665          670
351 Phe Lys Ile Asp Arg Ala Ser Glu Cys Arg Lys Lys Tyr Ala Phe Lys
352          675          680          685
354 Ala Cys His Pro Lys Ile Lys Ser Phe Tyr Phe Ala Ala Glu His Leu
355          690          695          700
357 Asp Asp Met Asn Arg Trp Leu Asn Arg Ile Asn Met Leu Thr Ala Gly
358 705          710          715          720
360 Tyr Ala Glu Arg Glu Arg Ile Lys Gln Glu Gln Asp Tyr Trp Ser Glu

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/508,343

DATE: 03/13/2005

TIME: 11:58:41

Input Set : A:\US10508343-seq list.txt

Output Set: N:\CRF4\03132005\J508343.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date